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<210> 5

<211> 908

<212> PRT

<213> Mus musculus

<400> 5

Met Pro Ser Ser Gly Pro Gly Asp Thr Ser Ser Ser Ser Leu Glu Arg

1

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Glu Asp Asp Arg Lys Glu Gly Glu Glu Gln Glu Glu Asn Arg Gly Lys

20

25

30

Glu Glu Arg Gln Glu Pro Ser Ala Thr Ala Arg Lys Val Gly Arg Pro

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35	40	45
Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Ser Asp Thr Pro Lys		
50	55	60
Asp Pro Ala Val Thr Thr Lys Ser Gln Pro Met Ala Gln Asp Ser Gly		
65	70	75
Pro Ser Asp Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg Ser Glu Pro		
85	90	95
Gln Pro Glu Glu Gly Ser Pro Ala Ala Gly Gln Lys Gly Gly Ala Pro		
100	105	110
Ala Glu Gly Glu Gly Thr Glu Thr Pro Pro Glu Ala Ser Arg Ala Val		
115	120	125
Glu Asn Gly Cys Cys Val Thr Lys Glu Gly Arg Gly Ala Ser Ala Gly		
130	135	140
Glu Gly Lys Glu Gln Lys Gln Thr Asn Ile Glu Ser Met Lys Met Glu		
145	150	155
Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp Glu Ser Ser Leu		
165	170	175
Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala Gly Asp Pro Tyr		
180	185	190
Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala Arg Trp Lys Arg		
195	200	205
Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Val Met Asn Ala Val Glu		
210	215	220
Glu Asn Gln Ala Ser Gly Glu Ser Gln Lys Val Glu Glu Ala Ser Pro		
225	230	235
Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro Thr Val Ala Thr		

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245	250	255
Thr Pro Glu Pro Val Gly Gly Asp Ala Gly Asp Lys Asn Ala Thr Lys		
260	265	270
Ala Ala Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg Gly Phe Gly Ile		
275	280	285
Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser Trp Trp Pro Gly		
290	295	300
Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg Ala Ala Glu Gly		
305	310	315
Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe Ser Val Val Cys		
325	330	335
Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser Ala Phe His Gln		
340	345	350
Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala Ile Tyr Glu Val		
355	360	365
Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe Pro Ala Cys His		
370	375	380
Asp Ser Asp Glu Ser Asp Ser Gly Lys Ala Val Glu Val Gln Asn Lys		
385	390	395
Gln Met Ile Glu Trp Ala Leu Gly Gly Phe Gln Pro Ser Gly Pro Lys		
405	410	415
Gly Leu Glu Pro Pro Glu Glu Glu Lys Asn Pro Tyr Lys Glu Val Tyr		
420	425	430
Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr Ala Pro Pro Pro		
435	440	445
Pro Ala Lys Lys Pro Arg Lys Ser Thr Thr Glu Lys Pro Lys Val Lys		

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450	455	460
Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val Tyr Glu Val Arg		
465	470	475 480
Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser Cys Gly Ser Leu		
	485	490 495
Asn Val Thr Leu Glu His Pro Leu Phe Ile Gly Gly Met Cys Gln Asn		
	500	505 510
Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr Asp Asp Asp Gly		
	515	520 525
Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg Glu Val Leu Met		
	530	535 540
Cys Gly Asn Asn Asn Cys Cys Arg Cys Phe Cys Val Glu Cys Val Asp		
545	550	555 560
Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile Lys Glu Asp Pro		
	565	570 575
Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr Gly Leu Leu Arg		
	580	585 590
Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe Phe Ala Asn Asn		
	595	600 605
His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro Pro Val Pro Ala		
	610	615 620
Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala		
625	630	635 640
Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln Val Asp Arg Tyr		
	645	650 655
Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val Gly Met Val Arg		

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660	665	670
His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg Ser Val Thr Gln		
675	680	685
Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser		
690	695	700
Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg Lys Gly Leu Tyr		
705	710	715 720
Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg Leu Leu His Asp		
725	730	735
Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe Trp Leu Phe Glu		
740	745	750
Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp Ile Ser Arg Phe		
755	760	765
Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu Val Ser Ala Ala		
770	775	780
His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro		
785	790	795 800
Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln Glu Cys Leu Glu		
805	810	815
His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr Ile Thr Thr Arg		
820	825	830
Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe Pro Val Phe Met		
835	840	845
Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met Glu Arg Val Phe		
850	855	860
Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met Ser Arg Leu Ala		

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Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val Pro Val Ile Arg His
885 890 895

Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys Val
900 905

<210> 6

<211> 859

<212> PRT

<213> Mus musculus

<400> 6

Met Lys Gly Asp Ser Arg His Leu Asn Glu Glu Glu Gly Ala Ser Gly
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Tyr Glu Glu Cys Ile Ile Val Asn Gly Asn Phe Ser Asp Gln Ser Ser
20 25 30

Asp Thr Lys Asp Ala Pro Ser Pro Pro Val Leu Glu Ala Ile Cys Thr
35 40 45

Glu Pro Val Cys Thr Pro Glu Thr Arg Gly Arg Arg Ser Ser Ser Arg
50 55 60

Leu Ser Lys Arg Glu Val Ser Ser Leu Leu Asn Tyr Thr Gln Asp Met
65 70 75 80

Thr Gly Asp Gly Asp Arg Asp Asp Glu Val Asp Asp Gly Asn Gly Ser
85 90 95

Asp Ile Leu Met Pro Lys Leu Thr Arg Glu Thr Lys Asp Thr Arg Thr
100 105 110

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg His Ser Asn Gly Thr Ser
115 120 125

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Ser Leu Glu Arg Gln Arg Ala Ser Pro Arg Ile Thr Arg Gly Arg Gln
130 135 140

Gly Arg His His Val Gln Glu Tyr Pro Val Glu Phe Pro Ala Thr Arg
145 150 155 160

Ser Arg Arg Arg Arg Ala Ser Ser Ser Ala Ser Thr Pro Trp Ser Ser
165 170 175

Pro Ala Ser Val Asp Phe Met Glu Glu Val Thr Pro Lys Ser Val Ser
180 185 190

Thr Pro Ser Val Asp Leu Ser Gln Asp Gly Asp Gln Glu Gly Met Asp
195 200 205

Thr Thr Gln Val Asp Ala Glu Ser Arg Asp Gly Asp Ser Thr Glu Tyr
210 215 220

Gln Asp Asp Lys Glu Phe Gly Ile Gly Asp Leu Val Trp Gly Lys Ile
225 230 235 240

Lys Gly Phe Ser Trp Trp Pro Ala Met Val Val Ser Trp Lys Ala Thr
245 250 255

Ser Lys Arg Gln Ala Met Pro Gly Met Arg Trp Val Gln Trp Phe Gly
260 265 270

Asp Gly Lys Phe Ser Glu Ile Ser Ala Asp Lys Leu Val Ala Leu Gly
275 280 285

Leu Phe Ser Gln His Phe Asn Leu Ala Thr Phe Asn Lys Leu Val Ser
290 295 300

Tyr Arg Lys Ala Met Tyr His Thr Leu Glu Lys Ala Arg Val Arg Ala
305 310 315 320

Gly Lys Thr Phe Ser Ser Ser Pro Gly Glu Ser Leu Glu Asp Gln Leu
325 330 335

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Lys Pro Met Leu Glu Trp Ala His Gly Gly Phe Lys Pro Thr Gly Ile
340 345 350

Glu Gly Leu Lys Pro Asn Lys Lys Gln Pro Val Val Asn Lys Ser Lys
355 360 365

Val Arg Arg Ser Asp Ser Arg Asn Leu Glu Pro Arg Arg Arg Glu Asn
370 375 380

Lys Ser Arg Arg Arg Thr Thr Asn Asp Ser Ala Ala Ser Glu Ser Pro
385 390 395 400

Pro Pro Lys Arg Leu Lys Thr Asn Ser Tyr Gly Gly Lys Asp Arg Gly
405 410 415

Glu Asp Glu Glu Ser Arg Glu Arg Met Ala Ser Glu Val Thr Asn Asn
420 425 430

Lys Gly Asn Leu Glu Asp Arg Cys Leu Ser Cys Gly Lys Lys Asn Pro
435 440 445

Val Ser Phe His Pro Leu Phe Glu Gly Gly Leu Cys Gln Ser Cys Arg
450 455 460

Asp Arg Phe Leu Glu Leu Phe Tyr Met Tyr Asp Glu Asp Gly Tyr Gln
465 470 475 480

Ser Tyr Cys Thr Val Cys Cys Glu Gly Arg Glu Leu Leu Leu Cys Ser
485 490 495

Asn Thr Ser Cys Cys Arg Cys Phe Cys Val Glu Cys Leu Glu Val Leu
500 505 510

Val Gly Ala Gly Thr Ala Glu Asp Ala Lys Leu Gln Glu Pro Trp Ser
515 520 525

Cys Tyr Met Cys Leu Pro Gln Arg Cys His Gly Val Leu Arg Arg Arg
530 535 540

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Lys Asp Trp Asn Met Arg Leu Gln Asp Phe Phe Thr Thr Asp Pro Asp
545 550 555 560

Leu Glu Glu Phe Glu Pro Pro Lys Leu Tyr Pro Ala Ile Pro Ala Ala
565 570 575

Lys Arg Arg Pro Ile Arg Val Leu Ser Leu Phe Asp Gly Ile Ala Thr
580 585 590

Gly Tyr Leu Val Leu Lys Glu Leu Gly Ile Lys Val Glu Lys Tyr Ile
595 600 605

Ala Ser Glu Val Cys Ala Glu Ser Ile Ala Val Gly Thr Val Lys His
610 615 620

Glu Gly Gln Ile Lys Tyr Val Asn Asp Val Arg Lys Ile Thr Lys Lys
625 630 635 640

Asn Ile Glu Glu Trp Gly Pro Phe Asp Leu Val Ile Gly Gly Ser Pro
645 650 655

Cys Asn Asp Leu Ser Asn Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu
660 665 670

Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr His Leu Leu Asn Tyr Thr
675 680 685

Arg Pro Lys Glu Gly Asp Asn Arg Pro Phe Phe Trp Met Phe Glu Asn
690 695 700

Val Val Ala Met Lys Val Asn Asp Lys Lys Asp Ile Ser Arg Phe Leu
705 710 715 720

Ala Cys Asn Pro Val Met Ile Asp Ala Ile Lys Val Ser Ala Ala His
725 730 735

Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly Met Asn Arg Pro Val
740 745 750

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Met Ala Ser Lys Asn Asp Lys Leu Glu Leu Gln Asp Cys Leu Glu Phe
755 760 765

Ser Arg Thr Ala Lys Leu Lys Lys Val Gln Thr Ile Thr Thr Lys Ser
770 775 780

Asn Ser Ile Arg Gln Gly Lys Asn Gln Leu Phe Pro Val Val Met Asn
785 790 795 800

Gly Lys Asp Asp Val Leu Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly
805 810 815

Phe Pro Ala His Tyr Thr Asp Val Ser Asn Met Gly Arg Gly Ala Arg
820 825 830

Gln Lys Leu Leu Gly Arg Ser Trp Ser Val Pro Val Ile Arg His Leu
835 840 845

Phe Ala Pro Leu Lys Asp Tyr Phe Ala Cys Glu
850 855

<210> 7

<211> 912

<212> PRT

<213> Homo sapiens

<400> 7

Met Pro Ala Met Pro Ser Ser Gly Pro Gly Asp Thr Ser Ser Ser Ala
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Ala Glu Arg Glu Glu Asp Arg Lys Asp Gly Glu Glu Gln Glu Glu Pro
20 25 30

Arg Gly Lys Glu Glu Arg Gln Glu Pro Ser Thr Thr Ala Arg Lys Val
35 40 45

Gly Arg Pro Gly Arg Lys Arg Lys His Pro Pro Val Glu Ser Gly Asp
50 55 60

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Thr Pro Lys Asp Pro Ala Val Ile Ser Lys Ser Pro Ser Met Ala Gln
65 70 75 80

Asp Ser Gly Ala Ser Glu Leu Leu Pro Asn Gly Asp Leu Glu Lys Arg
85 90 95

Ser Glu Pro Gln Pro Glu Glu Gly Ser Pro Ala Gly Gly Gln Lys Gly
100 105 110

Gly Ala Pro Ala Glu Gly Glu Gly Ala Ala Glu Thr Leu Pro Glu Ala
115 120 125

Ser Arg Ala Val Glu Asn Gly Cys Cys Thr Pro Lys Glu Gly Arg Gly
130 135 140

Ala Pro Ala Glu Ala Gly Lys Glu Gln Lys Glu Thr Asn Ile Glu Ser
145 150 155 160

Met Lys Met Glu Gly Ser Arg Gly Arg Leu Arg Gly Gly Leu Gly Trp
165 170 175

Glu Ser Ser Leu Arg Gln Arg Pro Met Pro Arg Leu Thr Phe Gln Ala
180 185 190

Gly Asp Pro Tyr Tyr Ile Ser Lys Arg Lys Arg Asp Glu Trp Leu Ala
195 200 205

Arg Trp Lys Arg Glu Ala Glu Lys Lys Ala Lys Val Ile Ala Gly Met
210 215 220

Asn Ala Val Glu Glu Asn Gln Gly Pro Gly Glu Ser Gln Lys Val Glu
225 230 235 240

Glu Ala Ser Pro Pro Ala Val Gln Gln Pro Thr Asp Pro Ala Ser Pro
245 250 255

Thr Val Ala Thr Thr Pro Glu Pro Val Gly Ser Asp Ala Gly Asp Lys
260 265 270

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Asn Ala Thr Lys Ala Gly Asp Asp Glu Pro Glu Tyr Glu Asp Gly Arg
275 280 285

Gly Phe Gly Ile Gly Glu Leu Val Trp Gly Lys Leu Arg Gly Phe Ser
290 295 300

Trp Trp Pro Gly Arg Ile Val Ser Trp Trp Met Thr Gly Arg Ser Arg
305 310 315 320

Ala Ala Glu Gly Thr Arg Trp Val Met Trp Phe Gly Asp Gly Lys Phe
325 330 335

Ser Val Val Cys Val Glu Lys Leu Met Pro Leu Ser Ser Phe Cys Ser
340 345 350

Ala Phe His Gln Ala Thr Tyr Asn Lys Gln Pro Met Tyr Arg Lys Ala
355 360 365

Ile Tyr Glu Val Leu Gln Val Ala Ser Ser Arg Ala Gly Lys Leu Phe
370 375 380

Pro Val Cys His Asp Ser Asp Glu Ser Asp Thr Ala Lys Ala Val Glu
385 390 395 400

Val Gln Asn Lys Pro Met Ile Glu Trp Ala Leu Gly Gly Phe Gln Pro
405 410 415

Ser Gly Pro Lys Gly Leu Glu Pro Pro Glu Glu Glu Lys Asn Pro Tyr
420 425 430

Lys Glu Val Tyr Thr Asp Met Trp Val Glu Pro Glu Ala Ala Ala Tyr
435 440 445

Ala Pro Pro Pro Pro Ala Lys Lys Pro Arg Lys Ser Thr Ala Glu Lys
450 455 460

Pro Lys Val Lys Glu Ile Ile Asp Glu Arg Thr Arg Glu Arg Leu Val
465 470 475 480

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Tyr Glu Val Arg Gln Lys Cys Arg Asn Ile Glu Asp Ile Cys Ile Ser
485 490 495

Cys Gly Ser Leu Asn Val Thr Leu Glu His Pro Leu Phe Val Gly Gly
500 505 510

Met Cys Gln Asn Cys Lys Asn Cys Phe Leu Glu Cys Ala Tyr Gln Tyr
515 520 525

Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Ile Cys Cys Gly Gly Arg
530 535 540

Glu Val Leu Met Cys Gly Asn Asn Asn Cys Cys Arg Cys Phe Cys Val
545 550 555 560

Glu Cys Val Asp Leu Leu Val Gly Pro Gly Ala Ala Gln Ala Ala Ile
565 570 575

Lys Glu Asp Pro Trp Asn Cys Tyr Met Cys Gly His Lys Gly Thr Tyr
580 585 590

Gly Leu Leu Arg Arg Arg Glu Asp Trp Pro Ser Arg Leu Gln Met Phe
595 600 605

Phe Ala Asn Asn His Asp Gln Glu Phe Asp Pro Pro Lys Val Tyr Pro
610 615 620

Pro Val Pro Ala Glu Lys Arg Lys Pro Ile Arg Val Leu Ser Leu Phe
625 630 635 640

Asp Gly Ile Ala Thr Gly Leu Leu Val Leu Lys Asp Leu Gly Ile Gln
645 650 655

Val Asp Arg Tyr Ile Ala Ser Glu Val Cys Glu Asp Ser Ile Thr Val
660 665 670

Gly Met Val Arg His Gln Gly Lys Ile Met Tyr Val Gly Asp Val Arg
675 680 685

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Ser Val Thr Gln Lys His Ile Gln Glu Trp Gly Pro Phe Asp Leu Val
690 695 700

Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Ile Val Asn Pro Ala Arg
705 710 715 720

Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe Phe Glu Phe Tyr Arg
725 730 735

Leu Leu His Asp Ala Arg Pro Lys Glu Gly Asp Asp Arg Pro Phe Phe
740 745 750

Trp Leu Phe Glu Asn Val Val Ala Met Gly Val Ser Asp Lys Arg Asp
755 760 765

Ile Ser Arg Phe Leu Glu Ser Asn Pro Val Met Ile Asp Ala Lys Glu
770 775 780

Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp Gly Asn Leu Pro Gly
785 790 795 800

Met Asn Arg Pro Leu Ala Ser Thr Val Asn Asp Lys Leu Glu Leu Gln
805 810 815

Glu Cys Leu Glu His Gly Arg Ile Ala Lys Phe Ser Lys Val Arg Thr
820 825 830

Ile Thr Thr Arg Ser Asn Ser Ile Lys Gln Gly Lys Asp Gln His Phe
835 840 845

Pro Val Phe Met Asn Glu Lys Glu Asp Ile Leu Trp Cys Thr Glu Met
850 855 860

Glu Arg Val Phe Gly Phe Pro Val His Tyr Thr Asp Val Ser Asn Met
865 870 875 880

Ser Arg Leu Ala Arg Gln Arg Leu Leu Gly Arg Ser Trp Ser Val Pro
885 890 895

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Val Ile Arg His Leu Phe Ala Pro Leu Lys Glu Tyr Phe Ala Cys Val
900 905 910

<210> 8

<211> 853

<212> PRT

<213> Homo sapiens

<400> 8

Met Lys Gly Asp Thr Arg His Leu Asn Gly Glu Glu Asp Ala Gly Gly
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Arg Glu Asp Ser Ile Leu Val Asn Gly Ala Cys Ser Asp Gln Ser Ser
20 25 30

Asp Ser Pro Pro Ile Leu Glu Ala Ile Arg Thr Pro Glu Ile Arg Gly
35 40 45

Arg Arg Ser Ser Ser Arg Leu Ser Lys Arg Glu Val Ser Ser Leu Leu
50 55 60

Ser Tyr Thr Gln Asp Leu Thr Gly Asp Gly Asp Gly Glu Asp Gly Asp
65 70 75 80

Gly Ser Asp Thr Pro Val Met Pro Lys Leu Phe Arg Glu Thr Arg Thr
85 90 95

Arg Ser Glu Ser Pro Ala Val Arg Thr Arg Asn Asn Asn Ser Val Ser
100 105 110

Ser Arg Glu Arg His Arg Pro Ser Pro Arg Ser Thr Arg Gly Arg Gln
115 120 125

Gly Arg Asn His Val Asp Glu Ser Pro Val Glu Phe Pro Ala Thr Arg

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130 135 140

Ser Leu Arg Arg Arg Ala Thr Ala Ser Ala Gly Thr Pro Trp Pro Ser
145 150 155 160

Pro Pro Ser Ser Tyr Leu Thr Ile Asp Leu Thr Asp Asp Thr Glu Asp
165 170 175

Thr His Gly Thr Pro Gln Ser Ser Ser Thr Pro Tyr Ala Arg Leu Ala
180 185 190

Gln Asp Ser Gln Gln Gly Gly Met Glu Ser Pro Gln Val Glu Ala Asp
195 200 205

Ser Gly Asp Gly Asp Ser Ser Glu Tyr Gln Asp Gly Lys Glu Phe Gly
210 215 220

Ile Gly Asp Leu Val Trp Gly Lys Ile Lys Gly Phe Ser Trp Trp Pro
225 230 235 240

Ala Met Val Val Ser Trp Lys Ala Thr Ser Lys Arg Gln Ala Met Ser
245 250 255

Gly Met Arg Trp Val Gln Trp Phe Gly Asp Gly Lys Phe Ser Glu Val
260 265 270

Ser Ala Asp Lys Leu Val Ala Leu Gly Leu Phe Ser Gln His Phe Asn
275 280 285

Leu Ala Thr Phe Asn Lys Leu Val Ser Tyr Arg Lys Ala Met Tyr His
290 295 300

Ala Leu Glu Lys Ala Arg Val Arg Ala Gly Lys Thr Phe Pro Ser Ser
305 310 315 320

Pro Gly Asp Ser Leu Glu Asp Gln Leu Lys Pro Met Leu Glu Trp Ala
325 330 335

His Gly Gly Phe Lys Pro Thr Gly Ile Glu Gly Leu Lys Pro Asn Asn

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340	345	350
Thr Gln Pro Val Val Asn Lys Ser Lys Val Arg Arg Ala Gly Ser Arg		
355	360	365
Lys Leu Glu Ser Arg Lys Tyr Glu Asn Lys Thr Arg Arg Arg Thr Ala		
370	375	380
Asp Asp Ser Ala Thr Ser Asp Tyr Cys Pro Ala Pro Lys Arg Leu Lys		
385	390	400
Thr Asn Cys Tyr Asn Asn Gly Lys Asp Arg Gly Asp Glu Asp Gln Ser		
405	410	415
Arg Glu Gln Met Ala Ser Asp Val Ala Asn Asn Lys Ser Ser Leu Glu		
420	425	430
Asp Gly Cys Leu Ser Cys Gly Arg Lys Asn Pro Val Ser Phe His Pro		
435	440	445
Leu Phe Glu Gly Gly Leu Cys Gln Thr Cys Arg Asp Arg Phe Leu Glu		
450	455	460
Leu Phe Tyr Met Tyr Asp Asp Asp Gly Tyr Gln Ser Tyr Cys Thr Val		
465	470	475
Cys Cys Glu Gly Arg Glu Leu Leu Leu Cys Ser Asn Thr Ser Cys Cys		
485	490	495
Arg Cys Phe Cys Val Glu Cys Leu Glu Val Leu Val Gly Thr Gly Thr		
500	505	510
Ala Ala Glu Ala Lys Leu Gln Glu Pro Trp Ser Cys Tyr Met Cys Leu		
515	520	525
Pro Gln Arg Cys His Gly Val Leu Arg Arg Arg Lys Asp Trp Asn Val		
530	535	540
Arg Leu Gln Ala Phe Phe Thr Ser Asp Thr Gly Leu Glu Tyr Glu Ala		

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545 550 555 560

Pro Lys Leu Tyr Pro Ala Ile Pro Ala Ala Arg Arg Arg Pro Ile Arg
 565 570 575

Val Leu Ser Leu Phe Asp Gly Ile Ala Thr Gly Tyr Leu Val Leu Lys
 580 585 590

Glu Leu Gly Ile Lys Val Gly Lys Tyr Val Ala Ser Glu Val Cys Glu
 595 600 605

Glu Ser Ile Ala Val Gly Thr Val Lys His Glu Gly Asn Ile Lys Tyr
 610 615 620

Val Asn Asp Val Arg Asn Ile Thr Lys Lys Asn Ile Glu Glu Trp Gly
625 630 635 640

Pro Phe Asp Leu Val Ile Gly Gly Ser Pro Cys Asn Asp Leu Ser Asn
 645 650 655

Val Asn Pro Ala Arg Lys Gly Leu Tyr Glu Gly Thr Gly Arg Leu Phe
 660 665 670

Phe Glu Phe Tyr His Leu Leu Asn Tyr Ser Arg Pro Lys Glu Gly Asp
 675 680 685

Asp Arg Pro Phe Phe Trp Met Phe Glu Asn Val Val Ala Met Lys Val
 690 695 700

Gly Asp Lys Arg Asp Ile Ser Arg Phe Leu Glu Cys Asn Pro Val Met
705 710 715 720

Ile Asp Ala Ile Lys Val Ser Ala Ala His Arg Ala Arg Tyr Phe Trp
 725 730 735

Gly Asn Leu Pro Gly Met Asn Arg Pro Val Ile Ala Ser Lys Asn Asp
 740 745 750

Lys Leu Glu Leu Gln Asp Cys Leu Glu Tyr Asn Arg Ile Ala Lys Leu

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TOP SECRET

755

760

765

Lys Lys Val Gln Thr Ile Thr Thr Lys Ser Asn Ser Ile Lys Gln Gly
770 775 780

Lys Asn Gln Leu Phe Pro Val Val Met Asn Gly Lys Glu Asp Val Leu
785 790 795 800

Trp Cys Thr Glu Leu Glu Arg Ile Phe Gly Phe Pro Val His Tyr Thr
805 810 815

Asp Val Ser Asn Met Gly Arg Gly Ala Arg Gln Lys Leu Leu Gly Arg
820 825 830

Ser Trp Ser Val Pro Val Ile Arg His Leu Phe Ala Pro Leu Lys Asp
835 840 845

Tyr Phe Ala Cys Glu
850

<210> 9

<211> 393

<212> DNA

<213> Mus musculus

<400> 9

tttctacagt atttcaggtg cctaccacac aggaacctt gaagaaaacc agtttctaga 60
agccgctgtt acctcttggt tacagtttat atatatatga tagatatgag atatatatat 120
ataaaaggta ctgttaacta ctgtacatcc cgacttcata atgggtgctt caaaacagcg 180
agatgagcaa agacatcagc ttccgcctgg ccctcgtgtg caaatggcgt ttcatgccca 240
tggatggtgt agaggggagc agctggaggg ggtttcacia actgaaggat gaccatatac 300
acccccacc cctgccccat gcctagcttc acctgccaaa aaggggctca gctgaggtgg 360
tcggaccctg gggaagctga gtgtggaatt tat 393

<210> 10

<211> 424

<212> DNA

<213> Mus musculus

09720086-074304

<400> 10

gaagaaaacc agtttctaga agccgctggt acctcttggt tacagtttat atatatatga 60
tagatatgag atatatatat ataaaaggta ctgttaacta ctgtacatcc cgacttcata 120
atggtgcttt caaacagcg agatgagcaa agacatcagc ttccgcctgg ccctctgtgc 180
aaagggtttc agcccaggat ggtgagaggg gagcatctgg aggggggttt aacaaactga 240
aggatgaccc atatcacccc ccacccctgc cccatgccta gcttcacctg ccaaaaagg 300
gctcagctga ggtggtcggg ccctggggaa gctgagtgtg gaatttatcc agactcgcgt 360
gcaataacct tagaatatga atctaaaatg actgcctcag aaaaatggct tgagaaaaca 420
ttgt 424

<210> 11

<211> 461

<212> DNA

<213> Mus musculus

<400> 11

tttaaagcaa accacagagg aggaaaacgc cggaggcttg gccttgcaaa agggttggac 60
atcatctcct gagttttcaa tgttaacctt cagtccatc taaaaagcaa aataggcccc 120
tccccttcgt tcccctccgg tcctaggagg cgaacttttt gttttctact ctttttcaga 180
gggggttttct gtttggttg gtttttggtt cttgctgtga ctgaaacaag agagttattg 240
cagcaaaatc agtaacaaca aaaagtagaa atgccttgga gcggaaaggg agagagggaa 300
aattctataa aaacttaaaa tattgggttt tttttttttc cttttctata tatctctttg 360
gttgtctcta gcctgatcag ataggagcac aaacaggaag agaatagaga ccctcggagg 420
cagagtctcc tctcccaccc cccgagcagt ctcaacagca c 461

<210> 12

<211> 465

<212> DNA

<213> Mus musculus

<400> 12

tcagaggggt tttctgttg tttgggtttt tgtttcttgc tgtgactgaa acaagagagt 60
tattgcagca aaatcagtaa caacaaaaag tagaaatgcc ttggagagga aaggagaga 120
gggaaaattc tataaaaact taaaatattg gttttttttt tttttccttt tctatatatc 180
tctttggttg tctctagcct gatcagatag gagcaciaaac aggaagagaa tagagaccct 240
cggaggcaga gtctcctctc ccaccccccg agcagtctca acagcaccat tcttggtcat 300
gcaaacaga acccaactag cagcagggcg ctgagagaac accacaccag acacttttct 360
acagtatttc aggtgcctac cacacaggaa accttgaaga aaaccagttt ctagaagccg 420

ctgttacctc ttgtttacag tttatatata tatgatagat atgag

465

<210> 13

<211> 393

<212> DNA

<213> Mus musculus

<400> 13

aaaacgccgg aggcctttgc cttgcacaag gggtggacat catctcctga gttttcaatg 60
ttaaccttca gtcctatcta aaaagcaaaa tagggccctc cccttcttcc cctccgggtcc 120
taggaggcga actttttgtt ttctactctt tttcagaggg gttttctgtt tgtttgggtt 180
tttgtttctt gctgtgactg aaacaagaga gttattgcag caaaatcagt aacaacaaaa 240
agtagaaatg ccttggagag gaaagggaga gagggaaaat tctataaaaa cttaaaatat 300
tggttttttt ttttttctt ttctatatat cgctttgggt gtctctagcc tgatcagata 360
ggagcacaaa caggaagaga atagagaccc tcg 393

<210> 14

<211> 309

<212> DNA

<213> Mus musculus

<400> 14

gtgatgattg acgccaaaga agtgtctgct gcacacaggg cccgttactt ctaggggtaa 60
ccttcctggc atgaacaggc ctttggatcc actgtgaatg ataagctgga gctgcaagag 120
tgtctggagc acggcagaat agccaagtgc agcaaagtga ggaccattac caccaggtca 180
aactctataa agcagggcaa agaccagcat ttccccgtct tcatgaacga gaaggaggac 240
atcctgtggg gcaactgaaat ggaaaggggc tttggcttcc ccgtccacta cacagacgtc 300
tccaacatg 309

<210> 15

<211> 341

<212> DNA

<213> Mus musculus

<400> 15

tgtaacctt cagtcctatc taaaaagcaa aataggcccc tccccttctt cccctccggg 60
cctaggaggc gaactttttg ttttctactc tttttcagag gggttttctg tttgtttggg 120
tttttgtttc ttgctgtgac tgaaacaaga gagttattgc agcaaaatca gtaacaacaa 180
aaagtagaaa tgccttggag aggaaagggg gagagggaaa attctataaa aacttaaaat 240

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attgggttttt ttttttttcc ttttttatat atctctttgg ttgtctctag cctgatcaga 300
taggagcaca aacaggaaga gaatagagac cctcggaggc a 341

<210> 16

<211> 240

<212> DNA

<213> Mus musculus

<220>

<221> Unsure

<222> (32)..(32)

<223> May be any nucleic acid

<400> 16

acatttttga tggttttttta ttgtctccag gngggggttaa tggcgggtca ctttccctca 60
ctctggaata tttctgatcc cacaaggggc cttcaacgtg gctgacgaat tcaaaatcag 120
ggacaatgtt ttctcaagcc atttttctga ggcagtcatt ttagattcat attctaaggt 180
tattgcacgc gagtctggat aaattccaca ctcagcttcc ccagggtccg accacctcag 240

<210> 17

<211> 256

<212> DNA

<213> Mus musculus

<220>

<221> Unsure

<222> (75)..(75)

<223> May be any nucleic acid

<400> 17

atcagcttcc gcctggccct ctgtgcaaag ggtttcagcc caggatgggg agaggggagc 60
agctggaggg ggttntaaca aactgaagga tgacccatat cccccccac cctgccccca 120
tgcttagctt cacctgcaa aaaggggctc agctgaggtg gtcggaccct ggggaagctg 180
agtgtggaat ttatccagac tcgcgtgcaa taaccttaga atatgaatct aaaatgactg 240
cctcagaaaa atggct 256

<210> 18

<211> 435

<212> DNA

<213> Mus musculus

<400> 18

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gtggaagccc atgcaatgat ctctctaacg tcaatcctgc ccgcaaaggt ttatatgagg 60
gcacaggaag gctcttcttc gagttttacc acttgctgaa ttatacccg cccaaggagg 120
gcgacaaccg tccattcttc tggatgttcg agaatgttgt ggccatgaaa gtgaatgaca 180
agaaagacat ctcaagattc ctggcatgta acccagtgat gatcgatgcc atcaagggtgt 240
ctgctgctca cagggcccgg tacttctggg gtaacctacc cggaatgaac aggcccgtga 300
tggcttcaaa gaatgataag ctcgagctgc aggactgcct ggagttcagt aggacagcaa 360
agttaaagaa agtgcagaca ataaccacca agtcgaactc catcagacag ggcaaaaacc 420
agcttttccc tgtag 435

<210> 19

<211> 522

<212> DNA

<213> Mus musculus

<400> 19

gatgatgtca gcagggatga catcaccacc tttagggctt ttccctggca ggggcccattg 60
tggctagtcc tcacgaagac tggagtagaa tgtttggagc tcaggaaggg tgggtggagt 120
ggagtctctt ccaggtgtga gggatacgaa ggaggaagct tagggaaatc cattccccac 180
tccctcttgc caaatgaggg gccagtgccc caacagctca ggtccccaga accccctagt 240
tcctcatgag aagctaggac cagaagcaca tcgttcccct tatctgagca gtgtttgggg 300
aactacagtg aaaaccttct ggagatgtta aaagcttttt accccacgat agattgtgtt 360
tttaaggggt gcttttttta ggggcatcac tggagataag aaagctgcat ttcagaaatg 420
ccatcgtaat ggttttttaa caccttttac ctaattacag gtgctatttt atagaagcag 480
acaacacttc tttttatgac tctcagactt ctattttcat gt 522

<210> 20

<211> 348

<212> DNA

<213> Mus musculus

<400> 20

aaaggaggcc cattagagtc ctgtctctgt ttgatggaat tgcaacgggg tacttggtgc 60
tcaaggagtt gggattataa gtggaaaagt acattgcctc cgaagtctgt gcagagtcca 120
tcgctgtggg aactgttaag catgaaggcc agatcaaata tgtcaatgac gtccggaaaa 180
tcaccaagaa aaatattgaa gagtggggcc cgttcgactt ggtgattggt ggaagcccat 240
gcaatgatct ctctaacgtc aatcctgccc gcaaagggtt atatgagggc acaggaaggc 300
tcttcttcga gttttaccac ttgctgaatt ataccgccc caaggagg 348

<210> 21

<211> 258
<212> DNA
<213> Mus musculus

<400> 21
gtttatggtt taagtcttcc tggcaccttc cccttgcttt ggtacaaggg ctgaagtcct 60
gttggctctg tagcatttcc caggatgatg atgtcagcag ggatgacatc atcaccttta 120
gggcttttcc ctggcagggg cccatgtggc tagtcctcac gaagactgga gtagaatgtt 180
tggagctcag gaaggggtggg tggagtgtgc ctcttccagg tgtgagggat acgaaggagg 240
aagcttaggg aaatccat 258

<210> 22
<211> 334
<212> DNA
<213> Mus musculus

<400> 22
tggggtaacc tacccggaat gaacagttaa agaaagtgca gacaataacc accaagtcga 60
actccatcag acagggcaaa aaccagcttt tccctgtagt catgaatggc aaggacgacg 120
ttttgtggtg cactgagctc gaaaggatct tcggcttccc tgctcactac acggacgtgt 180
ccaacatggg ccgcggcgcc cgtcagaagc tgctgggcag gtcctggagt gtaccgggtca 240
tcagacacct gtttgccccc ttgaaggact actttgcctg tgaatagttc taccaggac 300
tggggagctc tcggtcagag ccagtgccca gagt 334

<210> 23
<211> 299
<212> DNA
<213> Mus musculus

<220>
<221> Unsure
<222> (59)..(59)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (173)..(173)
<223> May be any nucleic acid

<400> 23
ctgtttttgt ttgttttttt ggtatcttag ccatcacttc tgagtgataa actcaggang 60

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gtaaaagaaa gccatcttac tacctacttc aagttttaa gtttcagggg aagagaacat 120
gagcaccatg ccgggctact ctaagcagcc aggtctgagc tgtgcacacg ganggagcac 180
cggggctccc ctgcaaggcc aggaggctct gctcccactg agcaggagaa agctgaggta 240
cagtgatgtg aggccccaca caggtgagct aaaaagggga caggtgaggt gccttcagg 299

<210> 24

<211> 455

<212> DNA

<213> Mus musculus

<400> 24

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gtgtctgtga gggccgtgaa ctgctgctgt gcagtaaacac aagctgctgc agatgcttct 120
gtgtggagtg tctggagggt ctggtgggag caggacagct gaggatgcca agctgcagga 180
accctggagc tgctatatgt gcctccctca gcgctgccat ggggtcctcc gacgcaggaa 240
agattggaac atgcgcctgc aagacttctt cactactgat cctgacctgg aagaatttca 300
ggagccaccc aagttgtacc cagcaattcc tgcagccaaa aggaggccca ttagagtcct 360
gtctctgttt gatggaattg caacggggta cttggtgctc aaggagttgg gtattaaagt 420
ggaaaagtac attgcctccg aagtctgtgc agagt 455

<210> 25

<211> 368

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (307)..(307)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (335)..(335)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (353)..(353)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (360)..(360)

<223> May be any nucleic acid

<400> 25

acgttttgta tgttttttta tttgctccag gtgggggttt gactgtcact ttcccacact 60
 ctggattagt tctgatccca ccacaaggag ccctcgaatt ggctaaagtg agaaactggg 120
 cctgaagact ccgtaccctc tgccatcttg ccgagggagt ctcttttag aaaacaatca 180
 aaggggtatt gcatgagtct ggatgaatcc cactctcagc ttgtccacgg gcccgaccac 240
 ctcatttagc cccctttttg gcaagggaga acctggctcc caagttctcc tccttcactt 300
 tcgttancaa accaaggggg aagaagccca ccgtngagaa cgcgccatct tgnaaagctn 360
 ggtcttcc 368

<210> 26

<211> 399

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (87)..(87)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (314)..(314)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (318)..(318)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (370)..(370)

<223> May be any nucleic acid

<400> 26

gaacatgagg atggagagaa gtatcagcac ccagaagaga aaaaggaatt taaaacaaaa 60
 accacagagg cggaaatacc ggaggcnttt gcttgcgaaa agggttggac atcatctcct 120
 gatTTTTcaa tgttattctt cagtcttatt taaaaacaaa accaagctcc cttcccttcc 180
 tcccccttcc ctttttttcc ggtcagacct tttattttct actcttttca gaggggtttt 240
 ctgtttgttt gggttttggt tcttgctgtg actgaaacaa gaagggttatt gcagcaaaaa 300
 tcaggtaaca aaanatantg aacaatacct tgcagaggaa aggtgggagg agaggaaaaa 360
 agggaaattn ctatagaaat ctatatattg gggttggtt 399

<210> 27

<211> 318

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<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (205)..(205)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (275)..(275)

<223> May be any nucleic acid

<400> 27

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caatgttacc ctggaacacc cctctcttctg tggaggaatg tgccaaaact gcaagaactg 120
ctttctggag tgtgctgacc agtacgacga cgacggctac cagtcctact gcaccatctg 180
ctgtggggggc cgtgaggtgc tcatntgcgg aaacaacaac tgctgcaggt gcttttgctg 240
ggagtgtgtg gacctcttgg tggggccggg ggctncccag gcagcagtta aggaagatca 300
tgtacgtcgg ggacgtcc                                     318
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<210> 28

<211> 259

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (227)..(227)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (234)..(234)

<223> May be any nucleic acid

<400> 28

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ggtgtgtgtt gagaagctga tgccgctgag ctggttttgc agtgcgttcc accaggccac 120
gtacaacaag cagcccatgt accgcaaagc catctacgag gtcctgcagg tggccagcag 180
ccgcgcgggg aagctgttcc cgggtgtgcca cgacagcgat gagagtnaca ctgncaaggc 240
cgtgggaggt gcagaacaa                                     259
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<210> 29

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<211> 483
<212> DNA
<213> Homo sapiens

<400> 29
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acactctgga ttagttctga tcccaccaca aggagccctc gaattggcta aagtgagaaa 120
ctggggcctga agactccgta cctcttgcca tcttgccgag ggagtctcct tttagaaaac 180
aatcaaaggg ttattgcatg agtctggatg aatcccactc tcagctgtcc acggggccga 240
ccacctcatc taggcccctt tttggcaagg agaaccgagg tcccaagttc tctcctttca 300
cttcgttaca aaccaggggg aaaaagccca cgtgaaaacg cggcatctgc aaaatggttc 360
cctttcttca tccctgggga aacctttgcg ccaaggcaac gtggaaactg atggttttac 420
tcaactcgct gttttgaagc gccattatga aatcgggggt gtacgtaggt aaagtcccg 480
gcc 483

<210> 30
<211> 337
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (41)..(41)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (45)..(45)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (176)..(176)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (190)..(190)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (207)..(207)
<223> May be any nucleic acid

<220>
<221> Unsure

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<222> (265)..(265)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (290)..(290)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (317)..(317)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (322)..(322)
<223> May be any nucleic acid

<400> 30
gggcattcag gtggaccgct acattgcctc ggaggtgtgt nagginctcca tcacgggtggg 60
catggtgagg caccagggga agatcatgta cgtcggggac gtccgcagcg tcacacagaa 120
gcatatccag gagggtgggc cattcgatct ggtgattggg ggcagtcctt gcaatnacct 180
ctccatcgtt aaccctgctc gcaaggncct ctacgagggc actggccggc tcttctttaa 240
gttctaccgc ctctgcatg atgcnegggc caaggagggg agatgatcgn cccttcttct 300
ggctctttaa gaatgtngtg gnccatgggc gtttagt 337

<210> 31
<211> 271
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (234)..(234)
<223> May be any nucleic acid

<400> 31
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actactgtac aaccgactt cataatgggtg ctttcaaaca gcgagatgag taaaaacatc 120
agcttccacg ttgccttctg cgcaaagggt ttcaccaagg atggagaaaagg ggagacagct 180
tgcagatggc gcgttctcac ggtgggctct tccccttggg ttgtaacgaa gtgnaggagg 240
agaacttggg agccagggtc tccctgccaa a 271

<210> 32
<211> 430

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<212> DNA

<213> Homo sapiens

<400> 32

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acgtttttgta tgtttttttta tttgctccag gtgggggtttt gactgtcact ttcccacact 60
ctggattagt tctgatccca ccacaaggag cctcgaatt ggctaaagtg agaaactggg 120
cctgaagact ccgtaccctc tgccatcttg ccgaggggagt ctcctttaga aaacaatcaa 180
agggttattg catgagtctg gatgaatccc actctcagct gtccacgggc ccgaccacct 240
catctagccc cctttttggc agggagaacc tggctcccaa gttctcctcc ttcacttcgt 300
tacaaaccaa ggggaagagc ccaccgtgag aacgcgccat ctgcaagctg tctccctttc 360
tccatccttg gtgaaacccc tttgcgcaga aggcaacgtg gaagctgatg tttttactca 420
tctcgtctgtt                                     430
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<210> 33

<211> 483

<212> DNA

<213> Homo sapiens

<400> 33

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ctggggcctga agactccgta ccctctgcca tcttgccgag ggagtctcct tttagaaaac 180
aatcaaaggg ttattgcatg agtctggatg aatcccactc tcagctgtcc acggggccga 240
ccacctcatc taggccccctt tttggcaagg agaaccggg tccaagttd tcctccttca 300
cttcgttaca aaccaggggg aaaaagccca cgtgaaaacg cggcatctgc aaaatggttc 360
cctttcttca tccttgggga aacctttgcg ccaaggcaac gtggaaactg atggttttac 420
tcaactcgct gttttgaagc gccattatga aatcgggggt gtacgtaggt aaagtcccgt 480
gcc                                     483
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<210> 34

<211> 411

<212> DNA

<213> Homo sapiens

<400> 34

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aaacaatcaa agggttattg catgagtctg gatgaatccc actctcagct gtccacgggc 240
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<212> DNA

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<400> 35

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gcttcccttt tttttcggtc agacctttta ttttctactc ttttcagagg ggttttctgt 420
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<210> 36

<211> 535

<212> DNA

<213> Homo sapiens

<400> 36

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tcgttacaaa ccacggggaa gagcccaccg tgagaacgcg ccactctgaa gctgtctccc 360
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aaggggttatt gcatgagtct ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240
tcatctagcc cccttttttg cagggagaac ctgggctccc aagttctcct ccttcacttc 300
gttacaaacc aaggggaagg agcccaccgt gagaacggcg ccatcttgca agctgtctcc 360
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tacagtattt caggtgccta ccacacagga aaccttgaag aaantcagtt tctaggaagc 240
cgctgttacc tcttgtttac agtttatata tatatgatag atatgagatn tatatataaa 300
agggtactgtt aactactgta caaccgcact tcataatggg tgctttcaaa caggcgaggt 360
gngtaaaaac atcagnttcc acgttngcct tttgcgcaaa gggtttcacc aggttgggga 420
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aaagggttat tgcattgagtc tggatgaatc ccactctcag ctgtccacgg gcccgaccac 240
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gcggcacatt tgccctccca gccactgagc tgctcgtgcc gcaccattcc tgggtcacgc 240
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gatcatgttg gagacgtcag tatagtggac tgggaaacca aatacccttt ccatttcagn 300
gcaccataag atgtcctctt tctcattcat gaagacaggg aaaatgctgg tctttggcct 360
gctcnatgga gtttgactcc gtagtaangg ccctcanttt ggntgacttg ggctatcctg 420
ncatgctcca gacacttccg nagggtcaca acagaagcat nttccagggg gtggngggcca 480
ttccgacctt tggnggattg ggggggaagc ccnaaaaat aacccttca aacggnnaaa 540
ccctngttcn gaangggccc cnttnccgang ggaaactggn ccgnttnttt ctttngggnt 600
tcctcccccc cccccnaaa ataatggng gccccagga ggggaattac cccccncn 660
ttnttttttt tttggaaatt tgggggccc ggggnnaann naaaanggn acttcnnnt 720
ttttggnccc nccnnnant ttnnncccaa aaanntaat taaaaaggcc cttttctggg 780
nccccnttn aaccgccccn ngatnggtnc ttggttccn aacacannnn cncaa 835

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<210> 52

<211> 479

<212> DNA

<213> Homo sapiens

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<220>
<221> Unsure
<222> (364)..(364)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (416)..(416)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (464)..(464)
<223> May be any nucleic acid

<400> 52
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gcctgaagac tccgtaccct ctgccatctt gccgagggag tctcctttta gaaaacaatc 180
aaagggttat tgcattgagtc tggatgaatc ccactctcag ctgtccacgg gcccgaccac 240
ctcatctagc cccctttttg gcagggagaa cctggctccc aagttctcct ccttcacttc 300
gttaciaaac aaggggaaga gccaccatg agaacgcgcc atctgcaagc tgtctccctt 360
tctncatcct tggtgaaacc tttgcgcaga aggcaacgtg gaagctgatg tttttntcat 420
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<210> 53
<211> 521
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (327)..(327)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (507)..(507)
<223> May be any nucleic acid

<400> 53
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tagatatgag atatatatat aaaagggtact gttaactact gtacaaccg acttcataat 180

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ggTgctttca aacagcgaga tgagtaaaaa catcagcttc cacgttgcct tctgcgcaaa 240
gggttttcacc aaggatggag aaagggagac agcttgacaga tggcgcgttc tcatgggtggg 300
ctcttccctt tggtttgtaa cgaagtntag gaggagaact tgggagccag gttctccctg 360
ccaaaaaggg ggctagatga ggtggtcggg cccgtggaca gctgagagtg ggattcatcc 420
agactcatgc aataaccctt tgattgtttc taaaaggaga ctccctcggc aagatggcag 480
agggtacgga gtcttcaggc ccagttntca ctttagccaa t 521

<210> 54

<211> 440

<212> DNA

<213> Homo sapiens

<400> 54

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aggggaagat catgtacgtc ggggacgtcc gcagcgtcac acagaagcat atccaggagt 180
ggggccatt cgatctgggtg attgggggca gtccctgcaa tgacctctcc atcgtcaacc 240
ctgctcgcga gggcctctac gagggcactg gccggctctt ctttgagttc taccgcctcc 300
tgcatgatgc gcggcccaag gagggagatg atcgccccctt cttctgggtc tttgagaatg 360
tggtggccat gggcgtttag tgacaagagg gacatctcgc gatttctcga gtccaaccct 420
gtgatgattg atgccaaaga 440

<210> 55

<211> 273

<212> DNA

<213> Homo sapiens

<400> 55

acgtttttgta tgtttttttta tttgctccag gtgggggtttt gactgtcact ttcccacact 60
ctggattagt tctgatccca ccacaaggag ccctcgaatt ggctaaagtg agaaactggg 120
cctgaagact ccgtaccctc tgccatcttg ccgagggagt ctcccttttag aaaacaatca 180
aagggttatt gcatgagtct ggatgaatcc cactctcagc tgtccacggg cccgaccacc 240
tcatctagcc cccttttttg cagggagaac ctg 273

<210> 56

<211> 190

<212> DNA

<213> Homo sapiens

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<220>
<221> Unsure
<222> (39)..(39)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (83)..(83)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (181)..(181)
<223> May be any nucleic acid

<400> 56
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agaagagaaa aaggaattta aancaaaaac cacagaggcg gaaataccgg agggctttgc 120
cttgcgaaaa gggttggaca tcctctctcg atttttcaat gttattcttc agtcctattt 180
naaaacaaag 190

<210> 57
<211> 445
<212> DNA
<213> Homo sapiens

<220>
<221> Unsure
<222> (167)..(167)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (353)..(353)
<223> May be any nucleic acid

<400> 57
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gcctaaaaat tagcaaaaac tgttgaaaca aggcacagtt ttttccccat atttgttacg 120
tcgtggctcc agttacaaaa aaattttaat gaaaacgtta aacatanaaa tagaagtttg 180
agattttaaa aagtgtataa aaagccccac aaaacttgtc aacgggttggt ccttattcta 240
caaaatagca ccagtaagaa gagtaaaagg tggtaaaaac catttatgac agcatttctg 300
aaatgcagct tgtctgaatt cccggttctc cctaaaaacg acttctttat ggnattaaaa 360
aagggtttta aaaaatctcc aaaggggagc accgagcttt gcagggttttc cctgtcatct 420
ctcagatgtg ggggaagctc gtggc 445

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<210> 58
 <211> 287
 <212> DNA
 <213> Homo sapiens

<220>
 <221> Unsure
 <222> (38)..(38)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (171)..(171)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (204)..(204)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (274)..(274)
 <223> May be any nucleic acid

<400> 58
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 ctgcatttca gaaatgctgt cataatggtt ttttaacacct tttactcctc nttactggtg 180
 ctatttttgt agaataaggg aacnacgttg acaagttttg gtgggggcct ttttatacac 240
 cttttttaaa atctccaact tcttaatttt taanggttta accgttt 287

<210> 59
 <211> 535
 <212> DNA
 <213> Homo sapiens

<220>
 <221> Unsure
 <222> (452)..(452)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (526)..(526)
 <223> May be any nucleic acid

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<400> 59

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cgtggctcca gttacaaaaa aatttttaatg aaaacgttaa acataaaaaat agaagtttga 180
gatttttaaaa agtgtataaa aagccccaca aaacttgatc acgttggtcc ttattctaca 240
aaatagcacc agtaagaaga gtaaaagggtg ttaaaaacca ttatgacagc atttctgaaa 300
tgcagcttgt ctgaattccc gttctcccta aaaacgactt cttatggaat aaaaaaggat 360
taaaaaatct ccaaaggag caccgagctt tgcagttttc cctgtccgtc tctcagatgt 420
ggggaaggta tgagaaatgt atgtctgtcc cngactgctg tcaactgcctc tgagttagta 480
aaaggtgaga atgagggtag cagcttccca tctggggcct gtgccngtgg aggggt 535

<210> 60

<211> 449

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (7)..(7)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (200)..(200)

<223> May be any nucleic acid

<400> 60

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tacgtgaacg acgtgaggaa catcaciaag aaaaatattg aagaatgggg cccatttgac 180
ttggtgattg gcggaaccan tgcaacgata tctcaaattg gaatccagcc aggaaaggcc 240
tgtatgaggg tacaggccgg ctcttcttcg aattttacca cctgctgaat tactcacgcc 300
ccaaggaggg tgatgaccgg ccgttcttct ggatgtttga gaatgttgta gccatgaagg 360
ttggcgacaa gagggacatc tcacggttcc tggagtgtaa tccagtgatg attgatgcca 420
tccaaagttt ctgctgctca caggggcccg 449

<210> 61

<211> 522

<212> DNA

<213> Homo sapiens

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<220>
 <221> Unsure
 <222> (146)..(146)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (281)..(281)
 <223> May be any nucleic acid

<220>
 <221> Unsure
 <222> (304)..(304)
 <223> May be any nucleic acid

<400> 61
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 atagcatcaa agaataataa actcngctg caggactgct tggaatacaa taggatagcc 180
 aagttaaaga agtacagac aataaccacc aagtcgaact cgatcaaaca ggggaaaaac 240
 caacttttcc ctgttgtcat gaatggcaaa gaagatgttt ngtggtgcac tgagctcgaa 300
 aggntctttg gctttcctgt gcactacaca gacgtgtcca acatgggccg tgggtgccgc 360
 cagaagctgc tgggaaggct ctggagcgtg cctgtcatcc gacacctctt cggccctctg 420
 aaggactact ttgcatgtga atagttccag ccaggggcca agcccactgg ggtgtgtggc 480
 agagcaggac ccaggaggtg tgattctgaa ggcaccccca gg 522

<210> 62
 <211> 573
 <212> DNA
 <213> Homo sapiens

<400> 62
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 tttacttaat cttgcttgat caggaactct ggtgtcttct tggcccccca cgtgatctcg 120
 ttcatgggtc cttttttgtt tatctcattt tctctgagggc tggctccttc tgttaacgtc 180
 ttggcatttg tgggaagcac aaaatgttct tgtccctcca actctgcttt tcgctccctg 240
 ccctgccatt cctctcccgc gcttgccctc tcccttccat ctttcccagg tacttttctc 300
 tcccagccct gccactcttc tgccgcacct gcgctctccc ctccatcttt ccaggtact 360
 tttgagcctt gactccccag gtcccttcat tctgtgctca ctccatgatg tcattttgtt 420
 ctccagttta agaaagtaca gacaataacc accaagtcga actcgatcaa acaggggaaa 480
 aaccaacttt tccctgttgt catgaatggc aaagaagatg ttttgtgggt cactgagctc 540
 gaaaggatct ttggctttcc tgtgcactac aca 573

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<210> 63
<211> 559
<212> DNA
<213> Homo sapiens

<400> 63
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ctaaaaatta gcaaaaactg ttgaaacaag gcacagtttt ttccccatat ttgttacgctc 120
gtggctccag ttacaaaaaa attttaaatga aaacggttaaa cataaaaaata gaagtttgag 180
attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttggttcct tattctacaa 240
aatagcacca gtaagaagag taaaagggtgt taaaaacccat tatgacagca tttctgaaat 300
gcagcttgctc tgaattcccg ttctccctaa aaacgacttc ttatggaata aaaaaggatt 360
aaaaaatctc caaaggggagc accgagcttt gcagttttcc ctgtcatcta tcagatgtgg 420
ggaagggtatg agaaatgtat gtctgtccct gactgctgtc actgcctctg agtttagtaa 480
aaagatgaga aatgagggtg gcagacttct catctgggga cctgtgcctg tggagggtag 540
gtctcctgga gaggggaatg 559

<210> 64
<211> 391
<212> DNA
<213> Homo sapiens

<400> 64
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tctttcagcc taaaaattag caaaaactgt tgaaacaagg cacagttttt tccccatatt 120
tgttacgtcg tggctccagt tacaaaaaaa attttaaatga aaacggttaaa cataaaaaata 180
gaagtttgag attttaaaaa gtgtataaaa agccccacaa aacttgtcaa cgttggttcct 240
tattctacaa aatagcacca gtaagaagag taaaagggtgt taaaaacccat tatgacagca 300
tttctgaaat gcagcttgctc tgaattcccg ttctccctaa aaacgacttc ttatggaata 360
aaaaaggatt aaaaaatctc caaaggggagc a 391

<210> 65
<211> 517
<212> DNA
<213> Homo sapiens

<400> 65
acaaatactg attttaatta aacataaggt aaactctagg caggggcatc tttcagccta 60

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aaaattagca aaaactgttg aaacaaggca cagttttttc cccatatttg ttacgtcgtg 120
gctccagtta cggaaaaatt ttaatgaaaa cgtaaacaat aaaaatagaa gtttgagatt 180
ttaaaaaagtg tataaaaaagc cccacaaaac ttgtcaacgt tgttccttat tctacaaaat 240
agcaccagta agaagagtaa aaggtgttaa aaaccattat gacagcattt ctgaaatgca 300
gcttgtctga attcccgttc tccctaaaaa cgacttctta tggaataaaa aaggattaaa 360
aaatctccaa agggagcacc gagctttgca gttttccctg tcatctctca gatgtgggga 420
aggatatgaga aatgtatgtc tgtccctgac tgctgtcact gcctctgagt ttagtaaaaa 480
gatgagaaat gagggtagca gacttctcat ctgggga 517

<210> 66

<211> 442

<212> DNA

<213> Homo sapiens

<400> 66

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taaaaattag caaaaactgt tgaaacaagg cacagttttt tccccatatt tgttacgtcg 120
tggtccagc tacaaaaaaa attttaatga aaacgttaaa cataaaaaata gaagtttgag 180
attttaaaaa gtgtataaaa agccccacaa aacttggtcaa cggtgttctt tattctacaa 240
aatagcacca gtaagaagag taaaaggtgt taaaaccat tatgacagca tttctgaaat 300
gcagcttgct tgaattcccg ttctccctaa aaacgacttc ttatggaata aaaaaggatt 360
aaaaaatctc caaaggagc accgagcttt gcagttttcc ctgtcatctc gcagatgtgg 420
ggaaggtatg agaaatgtat gt 442

<210> 67

<211> 396

<212> DNA

<213> Homo sapiens

<400> 67

gcagtcaggg acagacatac atttctcata ccttccccac atctgagaga tgacagggaa 60
aactgcaaag ctcggtgctc cctttggaga ttttttaatc cttttttttt ccataagaag 120
tcgttttttag ggagaacggg aattcagaca agctgcattt cagaaatgct gtcataatgg 180
tttttaacac cttttactct tcttactggg gctattttgt agaataagga acaacgttga 240
caagttttgt ggggcttttt atacactttt taaaatctca aacttctatt tttatgttta 300
acgttttcat taaaattttt ttgtaactgg agccacgacg taacaaatat ggggaaaaaa 360
ctgtgccttg tttcaacagt ttttgctaatt ttttag 396

<210> 68

<211> 287

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (7)..(7)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (169)..(169)

<223> May be any nucleic acid

<400> 68

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ctaaaaatta gcaaaaactg ttgaaacaag gcacagtttt tcccccatat ttgttacgtc 120
gtggctccag ttacaaaaaa aattttaatg aaaacgttaa acataaaaant agaagtttga 180
gatttttaaaa agtgtataaa aagccccaca aaacttggtca acgttggtcc ttattctaca 240
aaatagcacc agtaagaaga gtaaaagggtg ttaaaaacca ttatgac 287

<210> 69

<211> 356

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (193)..(193)

<223> May be any nucleic acid

<400> 69

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atgtgaatcc agccaggaaa ggcctgtatg aggggtacagg ccggctcttc ttcgaatttt 120
accacctgct gaattactca cgccccagg aggggtgatga ccggccgttc ttctggatgt 180
ttgagaatgt tgnagccatg aagggtggcg acaagaggga catctcacgg ttctggagt 240
gtaatccagt gatgattgat gccatcaaag tttctgctgc tcacagggcc cgatacttct 300
ggggcaacct acccgggatg aacaggatct ttggctttcc tgtgcactac acagac 356

<210> 70

<211> 408

<212> DNA

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<213> Homo sapiens

<220>

<221> Unsure

<222> (408)..(408)

<223> May be any nucleic acid

<400> 70

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agcctaataaa ttagcaaaaa ctgttgaaac aaggcacagt tttttcccca tatttgttac 120
gtcgtggctc cagttacaaa aaaaatttta atgaaaacgt taaacataaa aatagaagtt 180
tgagatttta aaaagtgtat aaaaagcccc acaaaaacttg tcaacgttgt tccttattct 240
acaaaatagc accagtaaga agagtaaaag gtgttaaaaa ccattatgac agcatttctg 300
aaatgcagct tgtctgaatt cccgttctcc ctaaaaacga cttcttatgg aataaaaaag 360
gattaaaaaa tctccaaagg gagcaccgag ctttgcagtt ttccctgn 408

<210> 71

<211> 439

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (50)..(50)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (85)..(85)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (405)..(405)

<223> May be any nucleic acid

<400> 71

gcattgtagct acaggacatt ttttaagggcc caggatcggt ttttcccagn tgcaagcaga 60
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ggagtctgca cgggacctat tagagtattt tccacaatga tgatgatttc agcaggggatg 180
acgtcatcat cacattcagg gctatttttt cccccacaaa cccaagggca gggggcactc 240
ttagctaaat ccctccccgt gactgcaata gaaccctctg gggagctcag gaaaggggggt 300
gtgctgagtt ctataatata agctgccata tattttgtag acaagtatgg ctctcccat 360

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atctccctct tccctaggag aggagtgtga aagcaaggga gcttngataa gacaccccct 420
caaaccatt ccctctcca 439

<210> 72

<211> 491

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (26)..(27)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (33)..(33)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (188)..(188)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (301)..(301)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (339)..(339)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (360)..(360)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (379)..(379)

<223> May be any nucleic acid

<400> 72

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aaaaaaaaatt ttaatgaaaa cgtaaacaat aaaaatagaa gtttgagatt ttaaaaaagt 180
tataaaaangc cccacaaaac ttgtcaacgt tgttccttat tctacaaaat agcaccagta 240
agaagagtaa aaggtgttaa aaaccattat gacagcattt ctgaaatgca gcttgtctga 300

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nttcccgttc tccctaaaaa cgactttotta tgggataana aagggattaa aaaatctccn 360
aaagggaggc accgagcttt gcaggttttc cctgggtcatc tctcaggatg tggggggagg 420
gtatggggaa atggtatggt ctggtccctg gactggctgg tcaactgcctc tgggggttng 480
gtaaaagggt g 491

<210> 73

<211> 443

<212> DNA

<213> Homo sapiens

<220>

<221> Unsure

<222> (9)..(9)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (11)..(11)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (23)..(24)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (126)..(126)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (157)..(157)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (170)..(170)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (341)..(341)

<223> May be any nucleic acid

<220>

<221> Unsure

<222> (347)..(347)

<223> May be any nucleic acid

<220>

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<221> Unsure
<222> (371)..(371)
<223> May be any nucleic acid

<220>
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<222> (405)..(405)
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<220>
<221> Unsure
<222> (412)..(412)
<223> May be any nucleic acid

<220>
<221> Unsure
<222> (430)..(430)
<223> May be any nucleic acid

<400> 73
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accggncgtt cttctggatg tttgagaatg ttgtagncat gaagggttggg gacaagaggg 180
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ctgtgcacta cacagacgtg tcccaacatg gggccgtggg ngccgcacca ggaagcttgc 360
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aaaaaaaaatt ttaatgaaaa cggttaaacad aaaaatagaa gtttgagatt ttaaaaagtg 180
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